



RAW SEQUENCE LISTING ERROR REPORT

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Application Serial Number: 09/553,431A
Source: 1600
Date Processed by STIC: 7/22/2003

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2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
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Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
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Revised 04/24/2003



1600

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/553,431A

DATE: 07/22/2003
 TIME: 13:26:19

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 5 <120> TITLE OF INVENTION: Manipulation of Min Genes in Plants
 7 <130> FILE REFERENCE: 920905.90041
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/553,431A
 C--> 10 <141> CURRENT FILING DATE: 2000-04-19
 12 <150> PRIOR APPLICATION NUMBER: 60/130,403
 13 <151> PRIOR FILING DATE: 1999-04-19
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 31 1 5 10 15
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 34 Pro Ser Ser Leu Ser Gln Lys Thr Leu Ile Ser Ser Pro Arg Phe Val
 35 20 25 30
 37 aat aac cct agc aga cgg agt cca ata cga tcc gtt ctt caa ttt aat 144
 38 Asn Asn Pro Ser Arg Arg Ser Pro Ile Arg Ser Val Leu Gln Phe Asn
 39 35 40 45
 41 cgc aaa ccg gaa ctc gcc gga aac acg ccg cgt atc gtc gtt atc acc 192
 42 Arg Lys Pro Glu Leu Ala Gly Glu Thr Pro Arg Ile Val Val Ile Thr
 43 50 55 60
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 51 85 90 95
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 57 aat tac act tgc gtc gag gtt ata aac gga gat tgt cgt ctc gat caa 384
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 59 115 120 125
 61 gct ctg gta cgt gat aag cgt tgg tcg aat ttc gaa ttg cta tgt ata 432
 62 Ala Leu Val Arg Asp Lys Arg Trp Ser Asn Phe Glu Leu Leu Cys Ile

Does Not Comply
Corrected Diskette Needed
pp 6-7

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216 Val Gly Leu Arg Asn Leu Asp Leu Leu Gly Leu Glu Asn Arg Val	
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223 gcc cta gtt cgt gat aaa cgc tgg tca aat ttc gaa ttg ctt tgt att	394
224 Ala Leu Val Arg Asp Lys Arg Trp Ser Asn Phe Glu Leu Leu Cys Ile	
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227 tca aaa cct agg tca aaa ttg cct tta gga ttt ggg gga aaa gct tta	442
228 Ser Lys Pro Arg Ser Lys Leu Pro Leu Gly Phe Gly Gly Lys Ala Leu	
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248 Ile Arg Asp Ile Lys Met Ile Val Asn Arg Val Arg Thr Asp Leu Ile	
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256 Leu Ser Leu Leu Ser Asp Thr Arg Gly Phe Glu Val Ile Arg Ser Thr	
257 230 235 240	
259 aat aga ggg ttt ccg ctt gtg ttg aac aag cct ccg act tta gca gga	826
260 Asn Arg Gly Phe Pro Leu Val Leu Asn Lys Pro Pro Thr Leu Ala Gly	
261 245 250 255	
263 ttg gca ttt gag cag gct gct tgg aga ttg gtt gag caa gat agc atg	874
264 Leu Ala Phe Glu Gln Ala Ala Trp Arg Leu Val Glu Gln Asp Ser Met	
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Input Set : A:\Un900411.app
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272 Phe Phe Gly Gly
273 295
275 tggtggagaa atgtgtcttg ttgttcatg taggagctgc tatgtgtcac ttgaaatgtt 1034
277 atgtgtacag taagctgata aggattgttt taattcagtt ttcaagagaga aaattagaat 1094
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297 35 40 45
299 Asn Arg Lys Pro Glu Leu Ala Gly Asp Thr Pro Arg Val Val Ala Ile
300 50 55 60
302 Asp Ala Asp Val Gly Leu Arg Asn Leu Asp Leu Leu Leu Gly Leu Glu
303 65 70 75 80
305 Asn Arg Val Asn Tyr Thr Val Val Glu Val Leu Asn Gly Asp Cys Arg
306 85 90 95
308 Leu Asp Gln Ala Leu Val Arg Asp Lys Arg Trp Ser Asn Phe Glu Leu
309 100 105 110
311 Leu Cys Ile Ser Lys Pro Arg Ser Lys Leu Pro Leu Gly Phe Gly Gly
312 115 120 125
314 Lys Ala Leu Val Trp Leu Asp Ala Leu Lys Asp Arg Gln Glu Gly Cys
315 130 135 140
317 Pro Asp Phe Ile Leu Ile Asp Cys Pro Ala Gly Ile Asp Ala Gly Phe
318 145 150 155 160
320 Ile Thr Ala Ile Thr Pro Ala Asn Glu Ala Val Leu Val Thr Thr Pro
321 165 170 175
323 Asp Ile Thr Ala Leu Arg Asp Ala Asp Arg Val Thr Gly Leu Leu Glu
324 180 185 190
326 Cys Asp Gly Ile Arg Asp Ile Lys Met Ile Val Asn Arg Val Arg Thr
327 195 200 205
329 Asp Leu Ile Arg Gly Glu Asp Met Met Ser Val Leu Asp Val Gln Glu
330 210 215 220
332 Met Leu Gly Leu Ser Leu Leu Ser Asp Thr Arg Gly Phe Glu Val Ile
333 225 230 235 240
335 Arg Ser Thr Asn Arg Gly Phe Pro Leu Val Leu Asn Lys Pro Pro Thr
336 245 250 255
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09/553,431A 6

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/553,431A

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Input Set : A:\Un900411.app
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Use of <220> Feature(NEW RULES):

Sequence(s) are missing the <220> Feature and associated headings.

Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence" or "Unknown". Please explain source of genetic material in <220> to <223> section (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp.29631-32) (Sec.1.823 of new Rules)

Seq#:5

VERIFICATION SUMMARY
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L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
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ORGANISM:Artificial Sequence
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